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1641
1653RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/910,009ADATE: 02/13/2002
TIME: 15:19:11Input Set : A:\2314-242rev.ST25.txt
Output Set: N:\CRF3\02132002\I910009A.raw

3 <110> APPLICANT: University of Utah Research Foundation
 4 Cognetix, Inc.
 5 Olivera, Baldomero M.
 6 McIntosh, J. Michael
 7 Garrett, James E.
 8 Watkins, Maren
 9 Cruz, Lourdes J.
 10 Shon, Ki-Joon
 11 Jacobsen, Richard
 12 Jones, Robert M.
 13 Cartier, G. Edward
 14 Shen, Greg S.
 15 Wagstaff, John D.

17 <120> TITLE OF INVENTION: Mu-Conopeptides
 19 <130> FILE REFERENCE: 2314-242

21 <140> CURRENT APPLICATION NUMBER: US/09/910,009A

21 <141> CURRENT FILING DATE: 2001-07-23

21 <150> PRIOR APPLICATION NUMBER: US 60/219,619

22 <151> PRIOR FILING DATE: 2000-07-21

24 <150> PRIOR APPLICATION NUMBER: US 60/245,157

25 <151> PRIOR FILING DATE: 2000-11-03

27 <150> PRIOR APPLICATION NUMBER: US 60/264,319

28 <151> PRIOR FILING DATE: 2001-01-29

30 <150> PRIOR APPLICATION NUMBER: US 60/277,270

31 <151> PRIOR FILING DATE: 2001-03-21

33 <160> NUMBER OF SEQ ID NOS: 520

35 <170> SOFTWARE: PatentIn version 3.0

37 <210> SEQ ID NO: 1

38 <211> LENGTH: 280

39 <212> TYPE: DNA

40 <213> ORGANISM: Conus arentus

42 <400> SEQUENCE: 1

43 caagaaggat	cgatagcagt	tcatgatgtc	taaactggga	gtcttcttga	ccatctgtat	60
45 gcttctgtt	cccttactg	ctttccgct	ggatggggat	caacctgcag	accgacctgc	120
47 agagcgtatg	caggacgact	ttataactga	gcatoatccc	ctgtttgatc	ctgtcaaacg	180
49 gtgttgcgag	aggccatgca	acataggatg	cgtacattgt	tgttaatgac	cagctttgtc	240
51 atcgcggcct	catcaagcga	ataagtaaaa	cgattgcagt			280

54 <210> SEQ ID NO: 2

55 <211> LENGTH: 67

56 <212> TYPE: PRT

57 <213> ORGANISM: Conus arentus

59 <400> SEQUENCE: 2

61 Met Met Ser Lys Leu Gly Val Phe Leu Thr Ile Cys Met Leu Leu Phe

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62 1 5 10 15
64 Pro Leu Thr Ala Leu Pro Leu Asp Gly Asp Gln Pro Ala Asp Arg Pro
65 20 25 30
67 Ala Glu Arg Met Gln Asp Asp Phe Ile Thr Glu His His Pro Leu Phe
68 35 40 45
70 Asp Pro Val Lys Arg Cys Cys Glu Arg Pro Cys Asn Ile Gly Cys Val
71 50 55 60
73 Pro Cys Cys
74 65
76 <210> SEQ ID NO: 3
77 <211> LENGTH: 14
78 <212> TYPE: PRT
79 <213> ORGANISM: Conus arentus
81 <220> FEATURE:
82 <221> NAME/KEY: PEPTIDE
83 <222> LOCATION: (1)..(14)
84 <223> OTHER INFORMATION: Xaa at residue 3 is Glu or gamma-carboxy Glu; Xaa at residue
5 an
85 d 12 is Pro or Hyp
88 <400> SEQUENCE: 3
90 Cys Cys Xaa Arg Xaa Cys Asn Ile Gly Cys Val Xaa Cys Cys 10
91 1 5
93 <210> SEQ ID NO: 4
94 <211> LENGTH: 244
95 <212> TYPE: DNA
96 <213> ORGANISM: Conus atlanticus
98 <400> SEQUENCE: 4
99 ggatccatga tgtctaaact gggagtcttg ttgaccatct gtctgcttct gtttccactt 60
101 actgctcttc cgctggatga agatcaaccg gtacaccgac ctgcagagcg tatgcaggac 120
103 atttcatctg atcaaçatct cttctttgat ctcataaac ggtgctgca gttgccatgc 180
105 gggccaggtt tttgcgtccc ttgttgctga catcaataac gtgttgatga ccaactttct 240
107 cgag 244
110 <210> SEQ ID NO: 5
111 <211> LENGTH: 69
112 <212> TYPE: PRT
113 <213> ORGANISM: Conus atlanticus
115 <400> SEQUENCE: 5
117 Gly Ser Met Met Ser Lys Leu Gly Val Leu Leu Thr Ile Cys Leu Leu 15
118 1 5 10 15
120 Leu Phe Pro Leu Thr Ala Leu Pro Leu Asp Glu Asp Gln Pro Val His 30
121 20 25 30
123 Arg Pro Ala Glu Arg Met Gln Asp Ile Ser Ser Asp Gln His Leu Phe 45
124 35 40 45
126 Phe Asp Leu Ile Lys Arg Cys Cys Glu Leu Pro Cys Gly Pro Gly Phe 60
127 50 55 60
129 Cys Val Pro Cys Cys
130 65
132 <210> SEQ ID NO: 6
133 <211> LENGTH: 15
134 <212> TYPE: PRT

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135 <213> ORGANISM: Conus atlanticus
137 <220> FEATURE:
138 <221> NAME/KEY: PEPTIDE
139 <222> LOCATION: (1)..(15)
140 <223> OTHER INFORMATION: Xaa at residue 3 is Glu or gamma-carboxy Glu; Xaa at residue

5, 8

141 and 13 is Pro or Hyp

144 <400> SEQUENCE: 6

W- 146 Cys Cys Xaa Leu Xaa Cys Gly Xaa Gly Phe Cys Val Xaa Cys Cys
147 1 5 10 15

149 <210> SEQ ID NO: 7

150 <211> LENGTH: 310

151 <212> TYPE: DNA

152 <213> ORGANISM: Conus aurisiacus

154 <400> SEQUENCE: 7

155 caagagggat cgatagcagt tcatgatgtc taaaactggga gtcttggtga ccatctgttt 60
156 gcttcgttt ccccttactg ctcttccgtat ggatggagat caatctgttag accgacactga 120
157 agagcgtatg caggacgaca tttcatctga gcagcatccc ttgtttaatc agaaaaagaat 180
158 gtgttgcggc gaaggccgga aatgccccag ctatttcaga aacagtcaaa tttgtcattg 240
159 ttgttaatgc acaacgtgtc gatgaccaac ttcgttatca cgactaatga ataagtaaaa 300
160 161 162 163 164 165

166 cgattgcagt

167 <210> SEQ ID NO: 8

168 <211> LENGTH: 74

169 <212> TYPE: PRT

170 <213> ORGANISM: Conus aurisiacus

171 <400> SEQUENCE: 8

172 Met Met Ser Lys Leu Gly Val Leu Leu Thr Ile Cys Leu Leu Phe 60
173 1 5 10 15
174 Pro Leu Thr Ala Leu Pro Met Asp Gly Asp Gln Ser Val Asp Arg Pro 120
175 20 25 30
176 Glu Glu Arg Met Gln Asp Asp Ile Ser Ser Glu Gln His Pro Leu Phe 180
177 35 40 45
178 Asn Gln Lys Arg Met Cys Cys Gly Glu Gly Arg Lys Cys Pro Ser Tyr 240
179 50 55 60
180 Phe Arg Asn Ser Gln Ile Cys His Cys Cys 300
181 65 70
182 183 184 185 186 187 188 189 190 191 192 193 194 195 196 197 198 199 200 201 202 203 204 205 206 207 208

190 <210> SEQ ID NO: 9

191 <211> LENGTH: 22

192 <212> TYPE: PRT

193 <213> ORGANISM: Conus aurisiacus

194 <220> FEATURE:

195 <221> NAME/KEY: PEPTIDE

196 <222> LOCATION: (1)..(22)
197 <223> OTHER INFORMATION: Xaa at residue 5 is Glu or gamma-carboxy Glu; Xaa at residue

10 i 198 s Pro or Hyp; Xaa at residue 12 is Tyr, 125I-Tyr, mono-iodo-Tyr,

199 di-iodo-Tyr, O-sulpho-Tyr or O-phospho-Tyr

200 201 202 203 <400> SEQUENCE: 9

W- 204 Met Cys Cys Gly Xaa Gly Arg Lys Cys Xaa Ser Xaa Phe Arg Asn Ser
205 1 5 10 15

206 Gln Ile Cys His Cys Cys

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Input Set : A:\2314-242rev.ST25.txt
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209 20
211 <210> SEQ ID NO: 10
212 <211> LENGTH: 257
213 <212> TYPE: DNA
214 <213> ORGANISM: Conus aurisiacus
216 <400> SEQUENCE: 10
217 ggatccatga tgtctaaact gggagtcttg ttgaccatct gtttgcttct gtttccccctt 60
219 actgctttc cgatcgatgg agatcaatct gtagaccgac ctgcagagcg tatgcaggat 120
221 gacatttcat ctgagcagca tcgcttggc aatcagaaaa gaaggtgctg ccggggcca 180
223 tgcccccggc aaatcgacgg tgaatattgt ggctgttgcc ttggatgata accgtgttga 240
225 tgaccaactt tctcgag 257
228 <210> SEQ ID NO: 11
229 <211> LENGTH: 75
230 <212> TYPE: PRT
231 <213> ORGANISM: Conus aurisiacus
233 <400> SEQUENCE: 11
235 Gly Ser Met Met Ser Lys Leu Gly Val Leu Leu Thr Ile Cys Leu Leu 15
236 1 5 10 15
238 Leu Phe Pro Leu Thr Ala Leu Pro Ile Asp Gly Asp Gln Ser Val Asp 30
239 20 25 30
241 Arg Pro Ala Glu Arg Met Gln Asp Asp Ile Ser Ser Glu Gln His Arg 45
242 35 40 45
244 Leu Phe Asn Gln Lys Arg Arg Cys Cys Arg Trp Pro Cys Pro Arg Gln 60
245 50 55 60
247 Ile Asp Gly Glu Tyr Cys Gly Cys Cys Leu Gly 75
248 65 70
250 <210> SEQ ID NO: 12
251 <211> LENGTH: 19
252 <212> TYPE: PRT
253 <213> ORGANISM: Conus aurisiacus
255 <220> FEATURE:
256 <221> NAME/KEY: PEPTIDE
257 <222> LOCATION: (1)..(19)
258 <223> OTHER INFORMATION: Xaa at residue 13 is Glu or gamma-carboxy Glu; Xaa at
residue 5 a
259 nd 7 is Pro or Hyp; Xaa at residue 4 is Trp or Bromo Trp; Xaa at 60
260 residue 14 is Tyr, 125I-Tyr, mono-iodo-Tyr, di-iodo-Tyr, O-sulpho
261 -Tyr or O-phospho-Tyr
264 <400> SEQUENCE: 12
266 Cys Cys Arg Xaa Xaa Cys Xaa Arg Gln Ile Asp Gly Xaa Xaa Cys Gly 15
267 1 5 10
269 Cys Cys Leu
272 <210> SEQ ID NO: 13
273 <211> LENGTH: 262
274 <212> TYPE: DNA
275 <213> ORGANISM: Conus aurisiacus
277 <400> SEQUENCE: 13
278 ggatccatga tgtctaaact gggagtcttg ttgaccatct gtctacttct gtttccccctt 60
280 actgctttc cgatggatgg agatcaacct gcagaccaac ctgcagatcg tatgcaggac 120
282 gacatttcat ctgagcagta tcccttggc gataagagac aaaagtgtt cactggaaag 180
283

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284 aaggggtcat gctccggcaa agcatgcaaa aatctcaa at gttgctctgg acgataacgt 240
286 gttgatgacc aacttctcg ag 262
289 <210> SEQ ID NO: 14
290 <211> LENGTH: 78
291 <212> TYPE: PRT
292 <213> ORGANISM: Conus aurisiacus
294 <400> SEQUENCE: 14
296 Gly Ser Met Met Ser Lys Leu Gly Val Leu Leu Thr Ile Cys Leu Leu
297 1 5 10 15
299 Leu Phe Pro Leu Thr Ala Phe Pro Met Asp Gly Asp Gln Pro Ala Asp
300 20 25 30
302 Gln Pro Ala Asp Arg Met Gln Asp Asp Ile Ser Ser Glu Gln Tyr Pro
303 35 40 45
305 Leu Phe Asp Lys Arg Gln Lys Cys Cys Thr Gly Lys Lys Gly Ser Cys
306 50 55 60
308 Ser Gly Lys Ala Cys Lys Asn Leu Lys Cys Cys Ser Gly Arg
309 65 70 75
311 <210> SEQ ID NO: 15
312 <211> LENGTH: 23
313 <212> TYPE: PRT
314 <213> ORGANISM: Conus aurisiacus
316 <220> FEATURE:
317 <221> NAME/KEY: PEPTIDE
318 <222> LOCATION: (1)..(23)
319 <223> OTHER INFORMATION: Xaa at residue 1 is Gln or pyro-Glu
322 <400> SEQUENCE: 15
324 Xaa Lys Cys Cys Thr Gly Lys Lys Gly Ser Cys Ser Gly Lys Ala Cys
W 325 1 5 10 15
327 Lys Asn Leu Lys Cys Cys Ser
328 20
330 <210> SEQ ID NO: 16
331 <211> LENGTH: 232
332 <212> TYPE: DNA
333 <213> ORGANISM: Conus aurisiacus
335 <400> SEQUENCE: 16
336 gatccatgta tgtctaaact gggagtcttg ctgaccatct gtctgcttct gtttccactt 60
338 actgctgttc cgctggatgg agatcaacct cttagaccgac acgcggagcg tatgcatgtat 120
340 ggcatttcac ctaaacgcca tccctggttt gatcccgta aacgggttg caaggtgcaa 180
342 tgcgagttt gcaccccttg ttgcttaacgt gttgatgacc aactttctcg ag 232
345 <210> SEQ ID NO: 17
346 <211> LENGTH: 68
347 <212> TYPE: PRT
348 <213> ORGANISM: Conus aurisiacus
350 <400> SEQUENCE: 17
352 Gly Ser Met Met Ser Lys Leu Gly Val Leu Leu Thr Ile Cys Leu Leu
353 1 5 10 15
355 Leu Phe Pro Leu Thr Ala Val Pro Leu Asp Gly Asp Gln Pro Leu Asp
356 20 25 30
358 Arg His Ala Glu Arg Met His Asp Gly Ile Ser Pro Lys Arg His Pro

Use of n and/or Xaa has been detected in the Sequence Listing.
Review the Sequence Listing to insure a corresponding
explanation is presented in the <220> to <223> fields of
each sequence using n or Xaa.

VERIFICATION SUMMARY
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Input Set : A:\2314-242rev.ST25.txt
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L:21 M:270 C: Current Application Number differs, Replaced Current Application No
L:21 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:90 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:146 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:205 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:266 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:324 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:381 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
L:438 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
L:494 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24
L:550 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:606 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30
L:609 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30
L:667 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33
L:727 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36
L:788 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39
L:851 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:42
L:911 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45
L:914 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:48
L:974 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:48
L:1037 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:1093 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:54
L:1152 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:57
L:1212 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:60
L:1215 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:60
L:1274 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:63
L:1277 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:63
L:1332 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66
L:1390 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:69
L:1447 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:72
L:1506 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:75
L:1567 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:78
L:1626 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:81
L:1685 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:84
L:1741 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:87
L:1800 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:90
L:1859 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:93
L:1915 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:96
L:1990 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:99
L:1993 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:99
L:2068 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:102
L:2071 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:102
L:2087 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:103
L:2090 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:103
L:2150 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:106
L:2209 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:109
L:2212 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:109
L:2274 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:112

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L:2332 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:115

L:2389 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:118

L:2392 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:118

L:2451 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:121